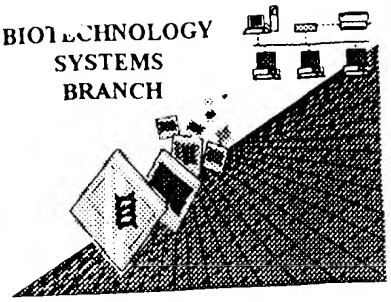

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/908,943

Source: OIPE

Date Processed by STIC: 7/30/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 3.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

http://www.uspto.gov/web/offices/pac/checker

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 09/908,943

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 ____ Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2 ____ Invalid Line Length	The rules require that a line **not exceed** 72 characters in length. This includes white spaces.
3 ____ Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.
4 ____ Non-ASCII	The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**
5 ____ Variable Length	Sequence(s) ____ contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6 ____ PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**
7 ____ Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.
8 ____ Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If **intentional**, please insert the following lines for **each** skipped sequence. <210> sequence id number <400> sequence id number 000
9 ____ Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.
10 ✓ Invalid <213> Response	Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence
11 ____ Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12 ____ PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13 ____ Misuse of n	n can only be used to represent a single **nucleotide** in a nucleic acid sequence. N is **not** used to represent any value not specifically a nucleotide.

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/908,943

DATE: 07/30/2001
TIME: 11:01:18

pp 6-7

Input Set : A:\00281.txt
Output Set: N:\CRF3\07302001\I908943.raw

<110> APPLICANT: Yan, Riqiang
Tomasselli, Alfredo G.
Gurney, Mark E.
Emmons, Thomas L.
Bienkowski, Mike J.
Heinrikson, Robert L.

<120> TITLE OF INVENTION: SUBSTRATES AND ASSAYS FOR BETA-SECRETASE ACTIVITY

<130> FILE REFERENCE: 29915/00281

C--> **<140> CURRENT APPLICATION NUMBER: US/09/908,943**

C--> **<141> CURRENT FILING DATE: 2001-07-19**

<160> NUMBER OF SEQ ID NOS: 197

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/908,943

DATE: 07/30/2001
TIME: 11:01:18

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Output Set: N:\CRF3\07302001\I908943.raw

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60 agaccaagct tgtttccctg ctggccaaag tcagtaggag aggatgcaca gtttgctatt 1980
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65 <211> LENGTH: 501
66 <212> TYPE: PRT
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74 20 25 30
76 Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu Thr Asp
77 35 40 45
79 Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val Glu Met Val
80 50 55 60
82 Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr
83 65 70 75 80
85 Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser
86 85 90 95
88 Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr
89 100 105 110
91 Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys Gly Val
92 115 120 125
94 Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp
95 130 135 140
97 Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala Asn Ile
98 145 150 155 160
100 Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser Asn Trp
101 165 170 175
103 Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Pro Asp Asp
104 180 185 190
106 Ser Leu Glu Pro Phe Phe Asp Ser Leu Val Lys Gln Thr His Val Pro
107 195 200 205
109 Asn Leu Phe Ser Leu His Leu Cys Gly Ala Gly Phe Pro Leu Asn Gln
110 210 215 220
112 Ser Glu Val Leu Ala Ser Val Gly Gly Ser Met Ile Ile Gly Gly Ile
113 225 230 235 240
115 Asp His Ser Leu Tyr Thr Gly Ser Leu Trp Tyr Thr Pro Ile Arg Arg
116 245 250 255
118 Glu Trp Tyr Tyr Glu Val Ile Ile Val Arg Val Glu Ile Asn Gly Gln
119 260 265 270
121 Asp Leu Lys Met Asp Cys Lys Glu Tyr Asn Tyr Asp Lys Ser Ile Val
122 275 280 285
124 Asp Ser Gly Thr Thr Asn Leu Arg Leu Pro Lys Lys Val Phe Glu Ala
125 290 295 300
127 Ala Val Lys Ser Ile Lys Ala Ala Ser Ser Thr Glu Lys Phe Pro Asp

RAW SEQUENCE LISTING DATE: 07/30/2001
PATENT APPLICATION: US/09/908,943 TIME: 11:01:18

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Output Set: N:\CRF3\07302001\I908943.raw

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131                 325                 330                 335
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134             340                 345                 350
136 Thr Asn Gln Ser Phe Arg Ile Thr Ile Leu Pro Gln Gln Tyr Leu Arg
137         355                 360                 365
139 Pro Val Glu Asp Val Ala Thr Ser Gln Asp Asp Cys Tyr Lys Phe Ala
140     370                 375                 380
142 Ile Ser Gln Ser Ser Thr Gly Thr Val Met Gly Ala Val Ile Met Glu
143 385                 390                 395                 400
145 Gly Phe Tyr Val Val Phe Asp Arg Ala Arg Lys Arg Ile Gly Phe Ala
146                 405                 410                 415
148 Val Ser Ala Cys His Val His Asp Glu Phe Arg Thr Ala Ala Val Glu
149             420                 425                 430
151 Gly Pro Phe Val Thr Leu Asp Met Glu Asp Cys Gly Tyr Asn Ile Pro
152         435                 440                 445
154 Gln Thr Asp Glu Ser Thr Leu Met Thr Ile Ala Tyr Val Met Ala Ala
155     450                 455                 460
157 Ile Cys Ala Leu Phe Met Leu Pro Leu Cys Leu Met Val Cys Gln Trp
158 465                 470                 475                 480
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161                 485                 490                 495
163 Ile Ser Leu Leu Lys
164             500
167 <210> SEQ ID NO: 3
168 <211> LENGTH: 1977
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170 <213> ORGANISM: Homo sapiens
172 <400> SEQUENCE: 3
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175 ctgcggctgc cccgggagac cgacgaagag cccgaggagc ccggccggag gggcagcttt 180
176 gtggagatgg tggacaacct gaggggcaag tcggggcagg gctactacgt ggagatgacc 240
177 gtgggcagcc ccccgcagac gctcaacatc ctggtggata caggcagcag taactttgca 300
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179 taccgggacc tccggaaggg tgtgtatgtg ccctacaccc agggcaagtg ggaaggggag 420
180 ctgggcaccg acctggtaag catcccccat ggccccaacg tcactgtgcg tgccaacatt 480
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184 acaggcagtc tctggtatac acccatccgg cgggagtggt attatgaggt gatcattgtg 720
185 cgggtggaga tcaatggaca ggatctgaaa atggactgca aggagtacaa ctatgacaag 780
186 agcattgtgg acagtggcac caccaacctt cgtttgccca agaaagtgtt tgaagctgca 840
187 gtcaaatcca tcaaggcagc ctcctccacg gagaagttcc ctgatggttt ctggctagga 900
188 gagcagctgg tgtgctggca agcaggcacc acccccttgga acattttccc agtcatctca 960
189 ctctacctaa tgggtgaggt taccaaccag tccttccgca tcaccatcct tccgcagcaa 1020
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191 tcacagtcat ccacgggcac tgttatggga gctgttatca tggagggctt ctacgttgtc 1140
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/908,943

DATE: 07/30/2001
TIME: 11:01:18

Input Set : A:\00281.txt
Output Set: N:\CRF3\07302001\I908943.raw

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194 aacattccac agacagatga gtcaaccctc atgaccatag cctatgtcat ggctgccatc 1320
195 tgcgccctct tcatgctgcc actctgcctc atggtgtgtc agtggcgctg cctccgctgc 1380
196 ctgcgccagc agcatgatga ctttgctgat gacatctccc tgctgaagtg aggaggccca 1440
197 tgggcagaag atagagattc ccctggacca cacctccgtg gttcactttg gtcacaagta 1500
198 ggagacacag atggcacctg tggccagagc acctcaggac cctccccacc caccaaatgc 1560
199 ctctgccttg atggagaagg aaaaggctgg caaggtgggt tccagggact gtacctgtag 1620
200 gaaacagaaa agagaagaaa gaagcactct gctggcggga atactcttgg tcacctcaaa 1680
201 tttaagtcgg gaaattctgc tgcttgaaac ttcagccctg aacctttgtc caccattcct 1740
202 ttaaattctc caacccaaag tattcttctt ttcttagttt cagaagtact ggcatcacac 1800
203 gcaggttacc ttggcgtgtg tccctgtggt accctggcag agaagagacc aagcttgttt 1860
204 ccctgctggc caaagtcagt aggagaggat gcacagtttg ctatttgctt tagagacagg 1920
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208 <211> LENGTH: 476
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217 20 25 30
219 Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu Thr Asp
220 35 40 45
222 Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val Glu Met Val
223 50 55 60
225 Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr
226 65 70 75 80
228 Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser
229 85 90 95
231 Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr
232 100 105 110
234 Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys Gly Val
235 115 120 125
237 Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp
238 130 135 140
240 Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala Asn Ile
241 145 150 155 160
243 Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser Asn Trp
244 165 170 175
246 Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Leu Cys Gly
247 180 185 190
249 Ala Gly Phe Pro Leu Asn Gln Ser Glu Val Leu Ala Ser Val Gly Gly
250 195 200 205
252 Ser Met Ile Ile Gly Gly Ile Asp His Ser Leu Tyr Thr Gly Ser Leu
253 210 215 220
255 Trp Tyr Thr Pro Ile Arg Arg Glu Trp Tyr Tyr Glu Val Ile Ile Val
256 225 230 235 240

RAW SEQUENCE LISTING DATE: 07/30/2001
PATENT APPLICATION: US/09/908,943 TIME: 11:01:18

Input Set : A:\00281.txt
Output Set: N:\CRF3\07302001\I908943.raw

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259                 245                 250                 255
261 Asn Tyr Asp Lys Ser Ile Val Asp Ser Gly Thr Thr Asn Leu Arg Leu
262             260                 265                 270
264 Pro Lys Lys Val Phe Glu Ala Ala Val Lys Ser Ile Lys Ala Ala Ser
265         275                 280                 285
267 Ser Thr Glu Lys Phe Pro Asp Gly Phe Trp Leu Gly Glu Gln Leu Val
268     290                 295                 300
270 Cys Trp Gln Ala Gly Thr Thr Pro Trp Asn Ile Phe Pro Val Ile Ser
271 305                 310                 315                 320
273 Leu Tyr Leu Met Gly Glu Val Thr Asn Gln Ser Phe Arg Ile Thr Ile
274                 325                 330                 335
277 Leu Pro Gln Gln Tyr Leu Arg Pro Val Glu Asp Val Ala Thr Ser Gln
278             340                 345                 350
280 Asp Asp Cys Tyr Lys Phe Ala Ile Ser Gln Ser Ser Thr Gly Thr Val
281         355                 360                 365
283 Met Gly Ala Val Ile Met Glu Gly Phe Tyr Val Val Phe Asp Arg Ala
284     370                 375                 380
286 Arg Lys Arg Ile Gly Phe Ala Val Ser Ala Cys His Val His Asp Glu
287 385                 390                 395                 400
289 Phe Arg Thr Ala Ala Val Glu Gly Pro Phe Val Thr Leu Asp Met Glu
290                 405                 410                 415
292 Asp Cys Gly Tyr Asn Ile Pro Gln Thr Asp Glu Ser Thr Leu Met Thr
293             420                 425                 430
295 Ile Ala Tyr Val Met Ala Ala Ile Cys Ala Leu Phe Met Leu Pro Leu
296         435                 440                 445
298 Cys Leu Met Val Cys Gln Trp Arg Cys Leu Arg Cys Leu Arg Gln Gln
299     450                 455                 460
301 His Asp Asp Phe Ala Asp Asp Ile Ser Leu Leu Lys
302 465                 470                 475
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306 <211> LENGTH: 14
307 <212> TYPE: PRT
308 <213> ORGANISM: Artificial Sequence
310 <220> FEATURE:
311 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic
312       peptide sequence
314 <400> SEQUENCE: 5
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316   1               5                  10
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320 <211> LENGTH: 15
321 <212> TYPE: PRT
322 <213> ORGANISM: Artificial Sequence
324 <220> FEATURE:
325 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic
326       peptide sequence
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09/908,943 -6

<210> SEQ ID NO 190
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<212> TYPE: PRT
<213> ORGANISM: synthetic peptide sequence

invalid - see item 10 on Error Summary Sheet

<400> SEQUENCE: 190
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1 5 10

09/908,943

<210> SEQ ID NO 194
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<212> TYPE: DNA
<213> ORGANISM: fusion protein comprising a maltose binding protein with 125 amino
acids from APP C-terminus.

nucleic sequence - also, this is a DNA sequence, not a PRT sequence

See Item 10 on Error Summary Sheet

FYI

Use of [illegible] has been detected in the Sequence Listing. [illegible] in the <220> to <223> fields of the [illegible] using n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: **US/09/908,943**

DATE: 07/30/2001
TIME: 11:01:19

Input Set : **A:\00281.txt**
Output Set: **N:\CRF3\07302001\I908943.raw**

L:15 M:270 C: Current Application Number differs, Replaced Current Application Number
L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:432 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:470 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:494 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:518 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:542 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:589 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:692 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:725 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:728 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:922 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41
L:1039 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49
L:1058 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50
L:1077 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:1096 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:1115 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53
L:1134 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54
L:1153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55
L:1172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:1191 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57
L:1210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58
L:1229 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:1248 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60
L:1267 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61
L:1286 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:62
L:1305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63
L:1324 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64
L:1343 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65
L:1362 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:1381 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67
L:1400 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68
L:1420 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69
L:1439 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70
L:1458 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:1477 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72
L:1496 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73
L:1515 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74
L:1534 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:75
L:1553 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:76
L:1572 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77
L:1591 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78
L:1610 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79
L:1629 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80
L:1653 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81
L:1677 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82
L:1701 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83

VERIFICATION SUMMARY
PATENT APPLICATION: **US/09/908,943**

DATE: 07/30/2001
TIME: 11:01:19

Input Set : **A:\00281.txt**
Output Set: **N:\CRF3\07302001\I908943.raw**

L:1725 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84
L:1749 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85
L:1773 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86
L:1792 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:87
L:4215 M:259 W: Allowed number of lines exceeded, <213> ORGANISM: